10/5█████

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Sequence Listing could not be accepted.
If you need help call the Patent Electronic Business Center at (866)
217-9197 (toll free).
Reviewer: Anne Corrigan
Timestamp: Thu Sep 13 17:19:26 EDT 2007
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Phe Ser Glu Pro Glu Ile Thr Leu Ile Ile Phe Gly Val Met Ala Gly
1   5   10   15

Val Ile Gly Thr Ile Leu Leu Ile Ser Tyr Gly Ile Arg Arg Leu Ile
            20   25   30

The above <222> (13)..(14) is erroneous for "D-Val"; Val is only at
location 13: "Met" is at location 14.

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<400> 25

Lys Lys Ile Thr Ala Gly Ala Ala Gly Val Ala Ala Gly Val Ala Ala
1   5   10   15

Ala

The above <223> response for the amino acid at location 6 is incorrect:
the <223> response shows "D-Glu." "Gly" is at location 6.

****

Application No: 10583996   Version No: 1.0

                    Input Set:

                    Output Set:

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                   Finished: 2007-09-04 12:49:57.061
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Input Set:

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       Assembly

<130> YEDA/038 PCT

<140> 10583996
<141> 2007-09-04

<150> US 60/530,899
<151> 2003-12-22

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<170> PatentIn version 3.3

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Glu Lys Asn Glu Gln Glu Leu Leu Glu Leu Asp Lys Trp Ala Ser Leu
            20   25   30

Trp Asn Trp Phe
        35

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Glu Lys Asn Glu Gln Glu Leu Leu Glu Leu Asp Lys Trp Ala Ser Leu
            20   25   30

Trp Asn Trp Phe
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1   5   10   15

Glu Lys Asn Glu Gln Glu Leu Leu Glu Leu Asp Lys Trp Ala Ser Leu
            20   25   30

Trp Asn Trp Phe
        35

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Ser Thr Met Gly Ala Arg Ser Met Thr Leu Thr Val Gln Ala Arg Gln
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Leu

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Leu

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Ser Thr Met Gly Ala Arg Ser Met Thr Leu Thr Val Gln Ala Arg Gln
            20   25   30

Leu

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            20   25   30

Leu

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1   5   10   15

Thr

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Thr Lys Lys

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1   5   10   15

Thr

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Thr Lys Lys

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Thr

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Thr Lys Lys

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            20   25   30

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Met Ala Gly Val Ile Gly Thr Ile Leu Leu Ile Ser Tyr Gly Ile Arg
            20   25   30

Arg Leu Ile
        35

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            20   25   30

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1   5   10   15

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1               5                   10                  15


Gly Ser Leu Lys Lys
            20


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<400>  22

Met Val Leu Gly Val Phe Ala Leu Leu Gln Leu Ile Ser Gly Ser Leu
1               5                   10                  15


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<400>  23

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1               5                   10                  15


Gly Ser Leu Lys Lys
            20


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1               5


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Lys Lys Ile Thr Ala Gly Ala Ala Gly Val Ala Ala Gly Val Ala Ala
1               5                   10                  15


Ala



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<400>  26

Trp Met Glu Trp Asp Arg Glu Ile Asn Asn Tyr Thr Ser Leu Ile His
1               5                   10                  15


Ser Leu Ile Glu Glu Ser Gln Asn Gln Gln Glu Lys Asn Glu Gln Glu
            20                  25                  30


Leu Leu



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<400> 27

Ala Val Gly Ile Gly Ala Leu Phe Leu Gly Phe Leu Gly
1   5   10

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<400> 28

Phe Phe Gly Ala Val Ile Gly Thr Ile Ala Leu Gly Val Ala Thr Ser
1   5   10   15

Ala Gln Ile Thr Ala Gly Ile Ala Leu Ala Glu Ala Arg Glu Ala Lys
            20   25   30

Arg

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<400> 29

Lys Lys Lys Met Val Leu Gly Val Phe Ala Leu Leu Phe Leu Ile Gly
1   5   10   15

Gly Ser Leu Lys Lys
            20